

290578US0XPCT.ST25.txt  
SEQUENCE LISTING

<110> Ishida, Nobuhiro  
Tokuhiro, Kenro  
Nagamori, Eiji  
Takahashi, Haruo  
Saito, Satoshi  
Ohni Shi, Tohru

<120> Promoter in the presence of organic acid and utilization thereof

<130> 290578US0XPCT

<140> 10/578,614  
<141> 2006-05-08

<150> PCT/JP04/16799  
<151> 2004-11-05

<150> JP 2003-379076  
<151> 2003-11-07

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<170> PatentIn version 3.3

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gcc gat ttg gca gaa agt ata atg aag aat ctt agg cggtg cat ccg Ala Asp Leu Ala Glu Ser Ile Met Lys Asn Leu Arg Arg Val His Pro 260 265 270			816
att tcc acc atg att aag ggt ctc tat gga ata aaa gag gat gtc ttc Ile Ser Thr Met Ile Lys Gly Leu Tyr Gly Ile Lys Glu Asp Val Phe 275 280 285			864
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Met Ala Cys Ala Ile Ser Ile Leu Met Lys Asp Leu Ala Asp Glu Val  
35 40 45

Ala Leu Val Asp Val Met Glu Asp Lys Leu Lys Gly Glu Met Met Asp  
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Leu Gln His Gly Ser Leu Phe Leu Arg Thr Pro Lys Ile Val Ser Gly  
65 70 75 80

Lys Asp Tyr Asn Val Thr Ala Asn Ser Arg Leu Val Ile Ile Thr Ala  
85 90 95

Gly Ala Arg Gln Gln Glu Gly Glu Ser Arg Leu Asn Leu Val Gln Arg

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Ser Gly Cys Asn Leu Asp Ser Ala Arg Phe Arg Tyr Leu Met Gly Glu  
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Arg Leu Gly Val His Pro Leu Ser Cys His Gly Trp Ile Leu Gly Glu  
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Val Ser Leu Lys Asn Leu His Pro Glu Leu Gly Thr Asp Ala Asp Lys  
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Ala Asp Leu Ala Glu Ser Ile Met Lys Asn Leu Arg Arg Val His Pro  
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	1 5 10		
	aaa gaa gaa cat gtt cca caa aat aaa att act att gtt ggt gtt ggt		99
	Lys Glu Glu His Val Pro Gln Asn Lys Ile Thr Ile Val Gly Val Gly		

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15	20	25				
gct gtt ggt atg gct tgt gct att tct att ttg atg aaa gat ttg gct Ala Val Gly Met Ala Cys Ala Ile Ser Ile Leu Met Lys Asp Leu Ala			147			
30	35	40	45			
gat gaa gtt gct ttg gtt gat gtt atg gaa gat aaa ttg aaa ggt gaa Asp Glu Val Ala Leu Val Asp Val Met Glu Asp Lys Leu Lys Gly Glu			195			
50	55		60			
atg atg gat ttg caa cat ggt tct ttg ttt ttg aga act cca aaa att Met Met Asp Leu Gln His Gly Ser Leu Phe Leu Arg Thr Pro Lys Ile			243			
65	70		75			
gtt tct ggt aaa gat tat aat gtt act gct aat tct aga ttg gtt att Val Ser Gly Lys Asp Tyr Asn Val Thr Ala Asn Ser Arg Leu Val Ile			291			
80	85		90			
att act gct ggt gct aga caa caa gaa ggt gaa tct aga ttg aat ttg Ile Thr Ala Gly Ala Arg Gln Gln Glu Gly Glu Ser Arg Leu Asn Leu			339			
95	100		105			
gtt caa aga aat gtt aat att ttt aaa ttt att att cca aat att gtt Val Gln Arg Asn Val Asn Ile Phe Lys Phe Ile Ile Pro Asn Ile Val			387			
110	115		120		125	
aaa tat tct cca aat tgt aaa ttg ttg gtt tct aat cca gtt gat Lys Tyr Ser Pro Asn Cys Lys Leu Leu Val Val Ser Asn Pro Val Asp			435			
130		135		140		
att ttg act tat gtt gct tgg aaa att tct ggt ttt cca aaa aat aga Ile Leu Thr Tyr Val Ala Trp Lys Ile Ser Gly Phe Pro Lys Asn Arg			483			
145		150		155		
gtt att ggt tct ggt tgt aat ttg gat tct gct aga ttt aga tat ttg Val Ile Gly Ser Gly Cys Asn Leu Asp Ser Ala Arg Phe Arg Tyr Leu			531			
160		165		170		
atg ggt gaa aga ttg ggt gtt cat cca ttg tct tgt cat ggt tgg att Met Gly Glu Arg Leu Gly Val His Pro Leu Ser Cys His Gly Trp Ile			579			
175		180		185		
ttg ggt gaa cat ggt gat tct tct gtt cca gtt tgg tct ggt gtt aat Leu Gly Glu His Gly Asp Ser Ser Val Pro Val Trp Ser Gly Val Asn			627			
190		195		200		205
gtt gct ggt gtt tct ttg aaa aat ttg cat cca gaa ttg ggt act gat Val Ala Gly Val Ser Leu Lys Asn Leu His Pro Glu Leu Gly Thr Asp			675			
210		215		220		
gct gat aaa gaa caa tgg aaa gct gtt cat aaa caa gtt gtt gat tct Ala Asp Lys Glu Gln Trp Lys Ala Val His Lys Gln Val Val Asp Ser			723			
225		230		235		
gct tat gaa gtt att aaa ttg aaa ggt tat act tct tgg gct att ggt Ala Tyr Glu Val Ile Lys Leu Lys Gly Tyr Thr Ser Trp Ala Ile Gly			771			
240		245		250		
ttg tct gtt gct gat ttg gct gaa tct att atg aaa aat ttg aga aga Leu Ser Val Ala Asp Leu Ala Glu Ser Ile Met Lys Asn Leu Arg Arg			819			
255		260		265		
gtt cat cca att tct act atg att aaa ggt ttg tat ggt att aaa gaa			867			

## 290578US0XPCT.ST25.txt

Val His Pro Ile Ser Thr Met Ile Lys Gly Leu Tyr Gly Ile Lys Glu	270 275 280 285	
gat gtt ttt ttg tct gtt cca tgt att ttg ggt caa aat ggt att tct	290 295 300	915
Asp Val Phe Leu Ser Val Pro Cys Ile Leu Gly Gln Asn Gly Ile Ser		
gat gtt gtt aaa gtt act ttg act cat gaa gaa gct tgt ttg aaa	305 310 315	963
Asp Val Val Lys Val Thr Leu Thr His Glu Glu Ala Cys Leu Lys		
aaa tct gct gat act ttg tgg ggt att caa aaa gaa ttg caa ttt taa	320 325 330	1011
Lys Ser Ala Asp Thr Leu Trp Gly Ile Gln Lys Glu Leu Gln Phe		
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His Val Pro Gln Asn Lys Ile Thr Ile Val Gly Val Gly Ala Val Gly	20 25 30	
Met Ala Cys Ala Ile Ser Ile Leu Met Lys Asp Leu Ala Asp Glu Val	35 40 45	
Ala Leu Val Asp Val Met Glu Asp Lys Leu Lys Gly Glu Met Met Asp	50 55 60	
Leu Gln His Gly Ser Leu Phe Leu Arg Thr Pro Lys Ile Val Ser Gly	65 70 75 80	
Lys Asp Tyr Asn Val Thr Ala Asn Ser Arg Leu Val Ile Ile Thr Ala	85 90 95	
Gly Ala Arg Gln Gln Glu Gly Glu Ser Arg Leu Asn Leu Val Gln Arg	100 105 110	
Asn Val Asn Ile Phe Lys Phe Ile Ile Pro Asn Ile Val Lys Tyr Ser	115 120 125	
Pro Asn Cys Lys Leu Leu Val Val Ser Asn Pro Val Asp Ile Leu Thr	130 135 140	

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Tyr Val Ala Trp Lys Ile Ser Gly Phe Pro Lys Asn Arg Val Ile Gly  
145 150 155 160

Ser Gly Cys Asn Leu Asp Ser Ala Arg Phe Arg Tyr Leu Met Gly Glu  
165 170 175

Arg Leu Gly Val His Pro Leu Ser Cys His Gly Trp Ile Leu Gly Glu  
180 185 190

His Gly Asp Ser Ser Val Pro Val Trp Ser Gly Val Asn Val Ala Gly  
195 200 205

Val Ser Leu Lys Asn Leu His Pro Glu Leu Gly Thr Asp Ala Asp Lys  
210 215 220

Glu Gln Trp Lys Ala Val His Lys Gln Val Val Asp Ser Ala Tyr Glu  
225 230 235 240

Val Ile Lys Leu Lys Gly Tyr Thr Ser Trp Ala Ile Gly Leu Ser Val  
245 250 255

Ala Asp Leu Ala Glu Ser Ile Met Lys Asn Leu Arg Arg Val His Pro  
260 265 270

Ile Ser Thr Met Ile Lys Gly Leu Tyr Gly Ile Lys Glu Asp Val Phe  
275 280 285

Leu Ser Val Pro Cys Ile Leu Gly Gln Asn Gly Ile Ser Asp Val Val  
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